C

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SEQUENCE LISTING

<110> Griffiths, Steven Gareth Ritchie, Rachel Jane Simard, Nathalie C.

<120> Hsp70 from Arthrobacter

<130> H-32534-A

<150> PCT/EP03/07602

<151> 2003-07-14

<150> GB 0216414.3

<151> 2002-07-15

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SEQLIST.TXT

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115 120 125 Ala Thr Lys Glu Ala Gly Glu Ile Ala Gly Leu Asn Val Leu Arg Ile 130 135 140 Val Asn Glu Pro Thr Ala Ala Ala Leu Ala Tyr Gly Leu Asp Lys Gly 145 150 155 160 Lys Glu Asp Glu Leu Ile Leu Val Phe Asp Leu Gly Gly Gly Thr Phe 165 170 175 Asp Val Ser Leu Leu Glu Val Gly Lys Asp Asp Asp Gly Phe Ser Thr 180 185 190 Ile Gln Val Arg Ala Thr Ser Gly Asp Asn Arg Leu Gly Gly Asp Asp 195 200 205 Trp Asp Gln Arg Ile Val Asp Tyr Leu Leu Asn Gln Leu Lys Val Lys 210 220 Gly Ile Asp Leu Ser Lys Asp Lys Ile Ala Leu Gln Arg Leu Arg Glu 225 230 235 240 Ala Ser Glu Gln Ala Lys Lys Glu Leu Ser Ser Ala Thr Ser Thr Asn 245 250 255 Ile Ser Leu Gln Tyr Leu Ser Val Thr Pro Asp Gly Pro Val His Leu 260 265 270 Asp Glu Gln Leu Thr Arg Ala Lys Phe Gln Glu Leu Thr Ala Asp Leu 275 280 285 Leu Glu Arg Thr Lys Lys Pro Phe Gln Asp Val Ile Ala Glu Ala Gly 290 295 300 Ile Lys Val Ser Asp Ile Asp His Ile Val Leu Val Gly Gly Ser Thr 305 310 315 320 Arg Met Pro Ala Val Thr Glu Leu Val Lys Gln Leu Ala Gly Gly Lys 325 330 335 Glu Pro Asn Lys Gly Val Asn Pro Asp Glu Val Val Ala Val Gly Ala 340 345 350 Ala Leu Gln Ala Gly Val Leu Lys Gly Glu Arg Lys Asp Val Leu Leu 355 360 365

SEQLIST.TXT Ile Asp Val Thr Pro Leu Ser Leu Gly Île Glu Thr Lys Gly Gly Val 370 375 380 Met Thr Lys Leu Ile Glu Arg Asn Thr Ala Ile Pro Thr Lys Arg Ser 385 390 395 400 Glu Thr Phe Thr Thr Ala Asp Asp Asn Gln Pro Ser Val Ala Ile Gln 405 410 415Val Phe Gln Gly Glu Arg Glu Phe Thr Arg Asp Asn Lys Pro Leu Gly 420 425 430 Thr Phe Glu Leu Thr Gly Ile Ala Pro Ala Pro Arg Gly Val Pro Gln
435 440 445 Val Glu Val Thr Phe Asp Ile Asp Ala Asn Gly Ile Val His Val Ser 450 460 Ala Lys Asp Lys Gly Thr Gly Lys Glu Gln Ser Met Thr Ile Thr Gly 465 470 475 480 Gly Ser Ser Leu Ser Lys Glu Asp Ile Glu Arg Met Val Ala Asp Ala 485 490 495 Glu Ala His Ala Ala Glu Asp Lys Ala Arg Arg Glu Gln Ala Glu Ala 500 505 510 Arg Asn Ser Ala Glu Gln Leu Ala Tyr Ser Val Asp Lys Ile Leu Thr 515 520 525 Asp Asn Asp Asp Lys Leu Pro Glu Glu Val Lys Thr Glu Val Lys Ala 530 540 Asp Val Gly Ala Leu Lys Thr Ala Leu Ala Gly Thr Asp Glu Asp Ala 545 550 555 560 Val Glu Ála Ala Ser Glu Lys Leu Gln Ala Ser Gln Thr Lys Leu Gly
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20